

009790" TTT9550

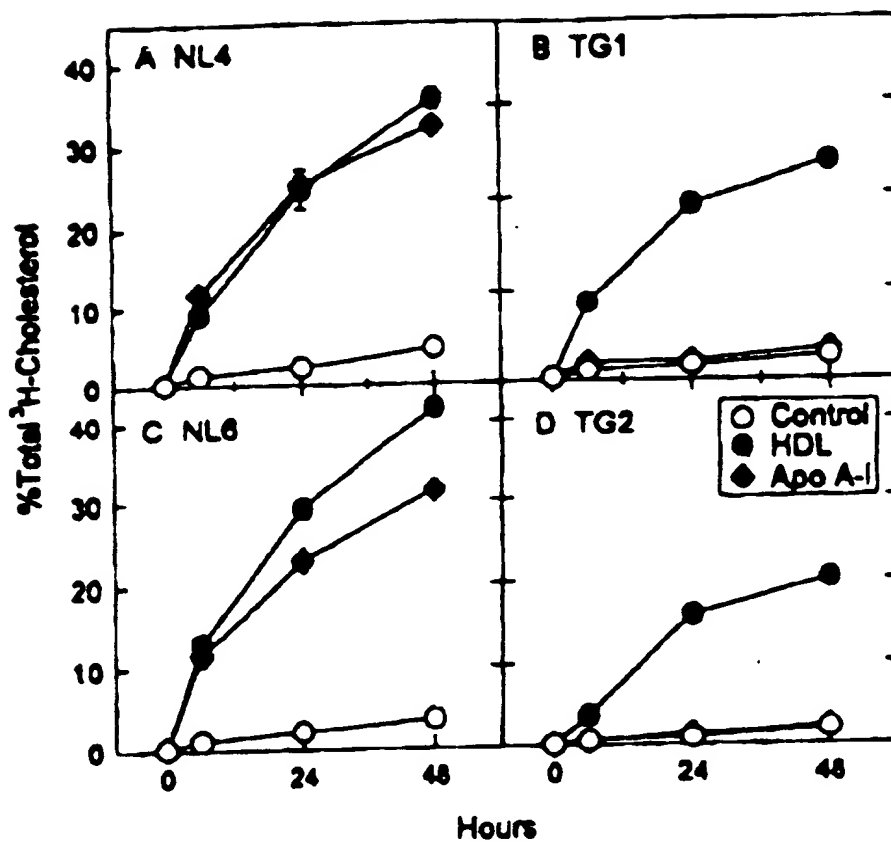


FIG. 1

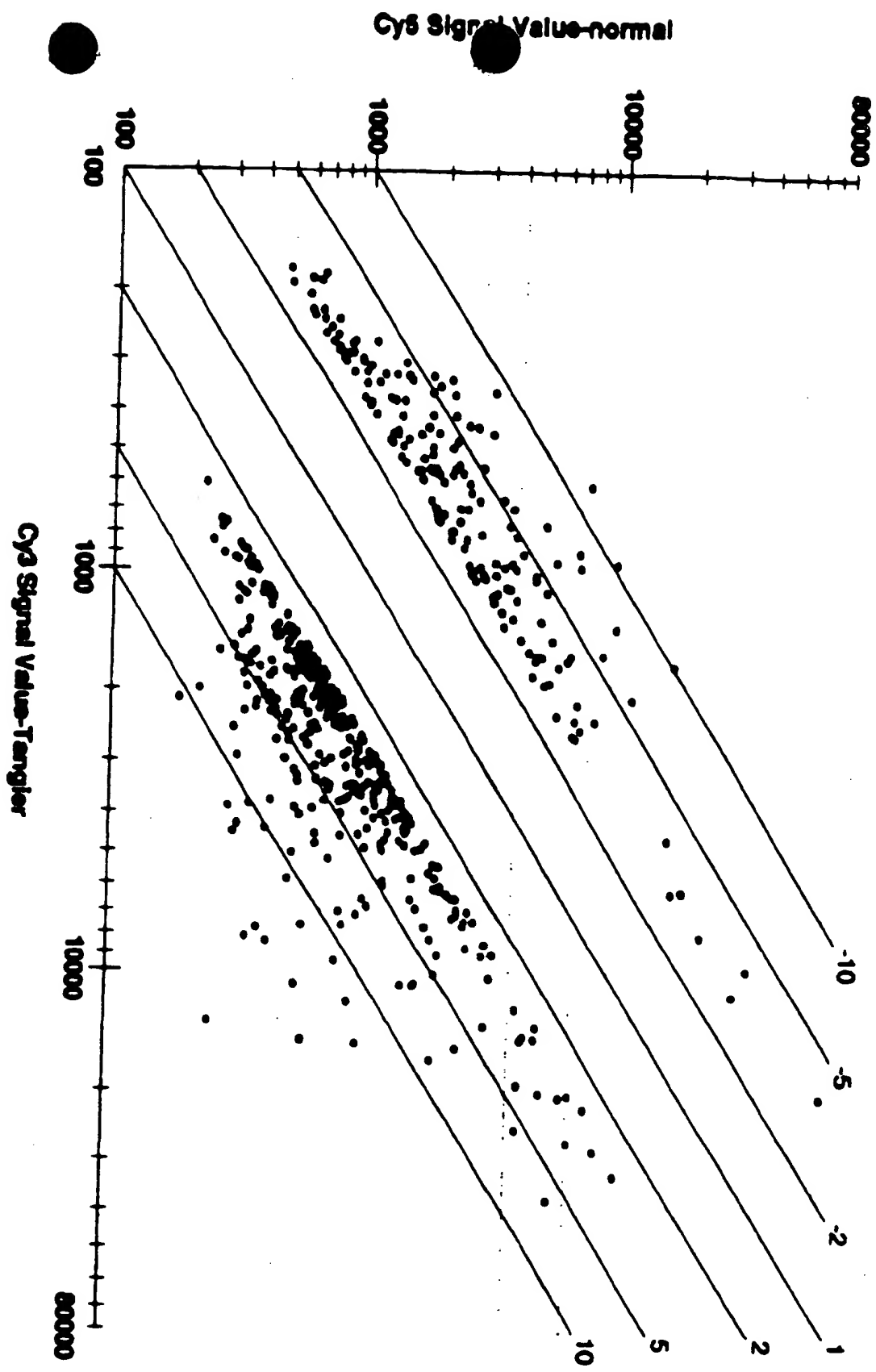


FIG. 2

09596441-061600

00596141.061600

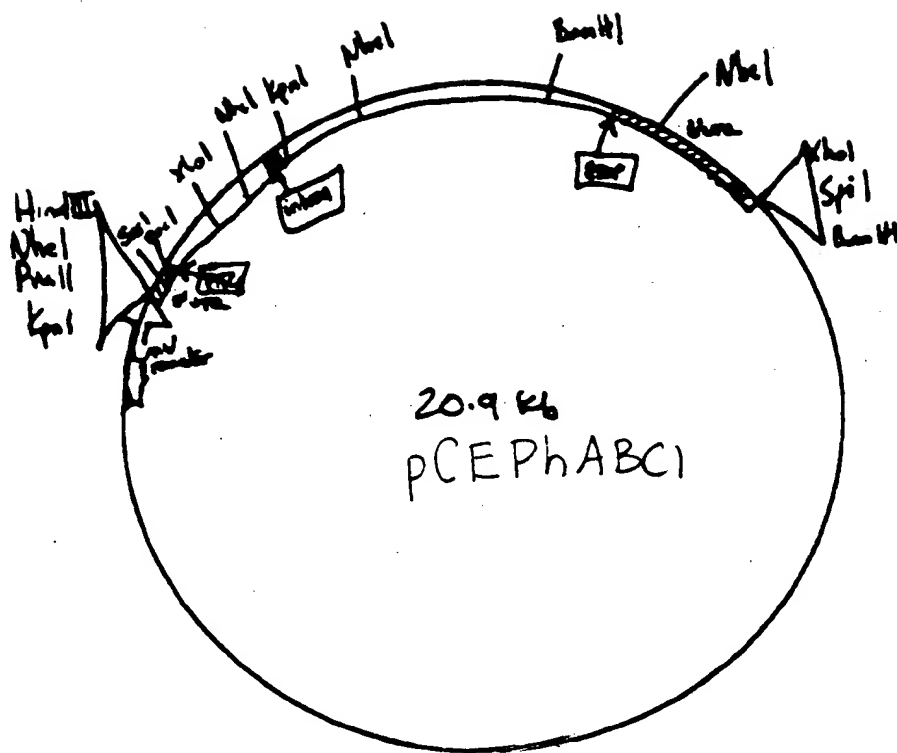


FIG. 3

# CVT ABCA1 Gene Structure

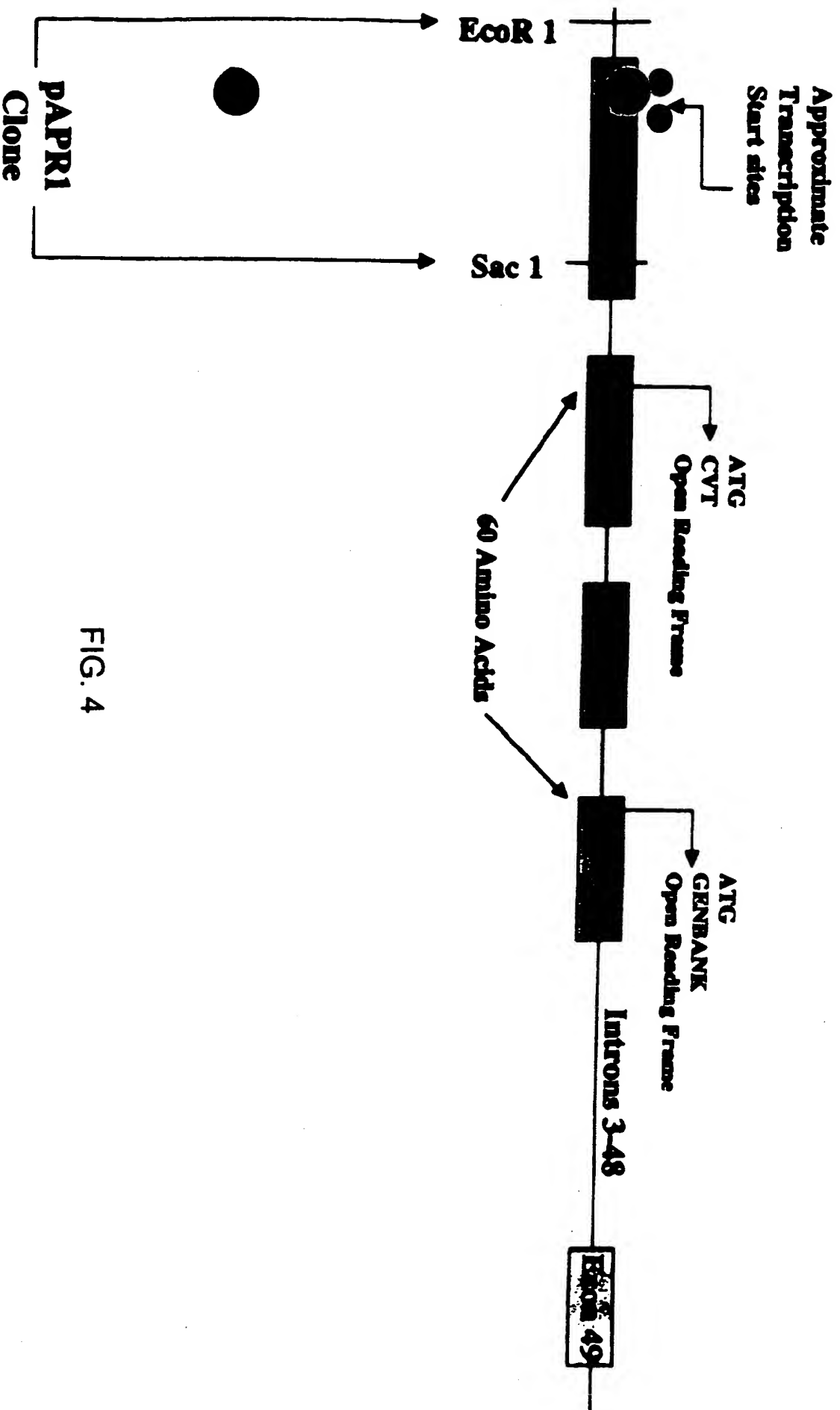


FIG. 4

apoA-I specific cholesterol efflux

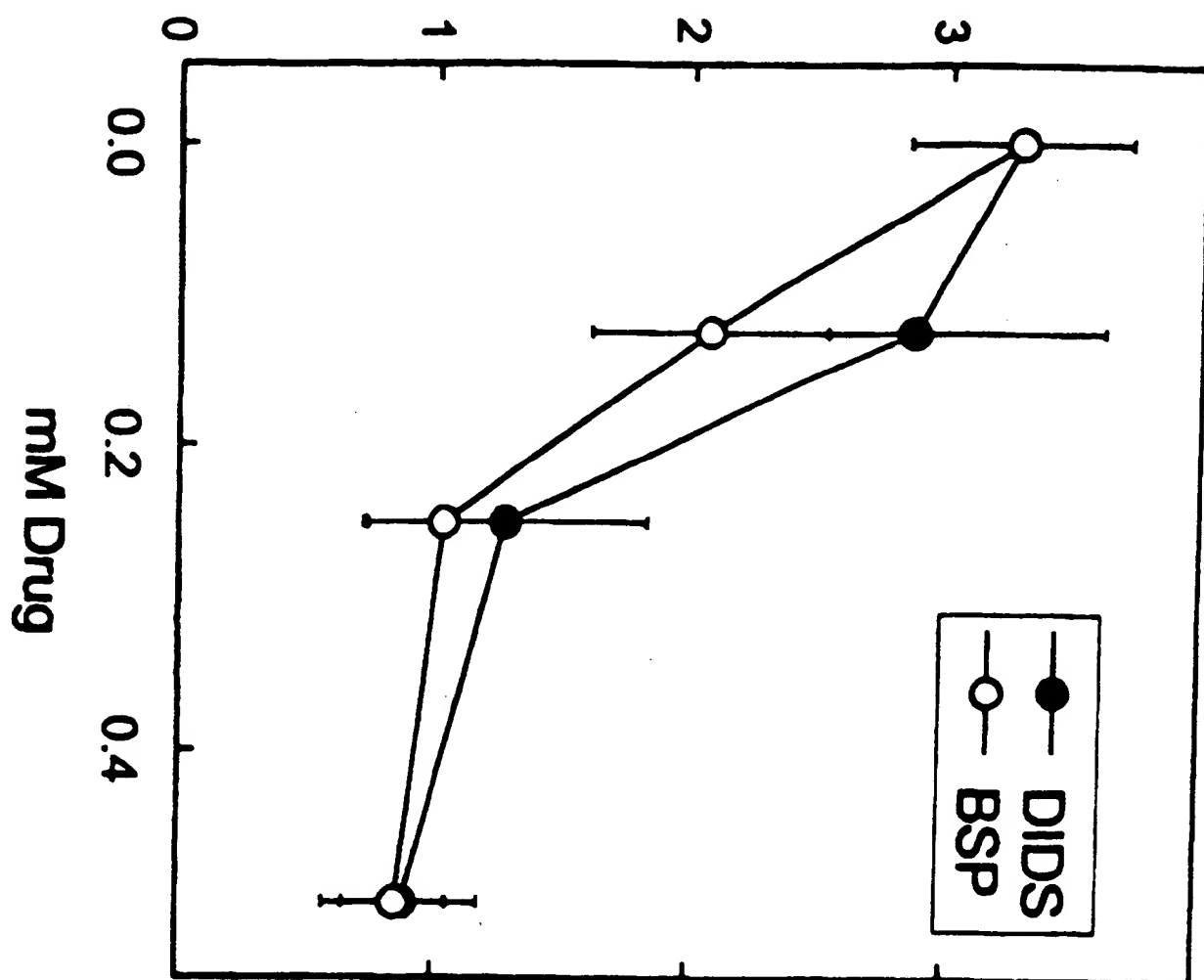


FIG. 5 95141-051500

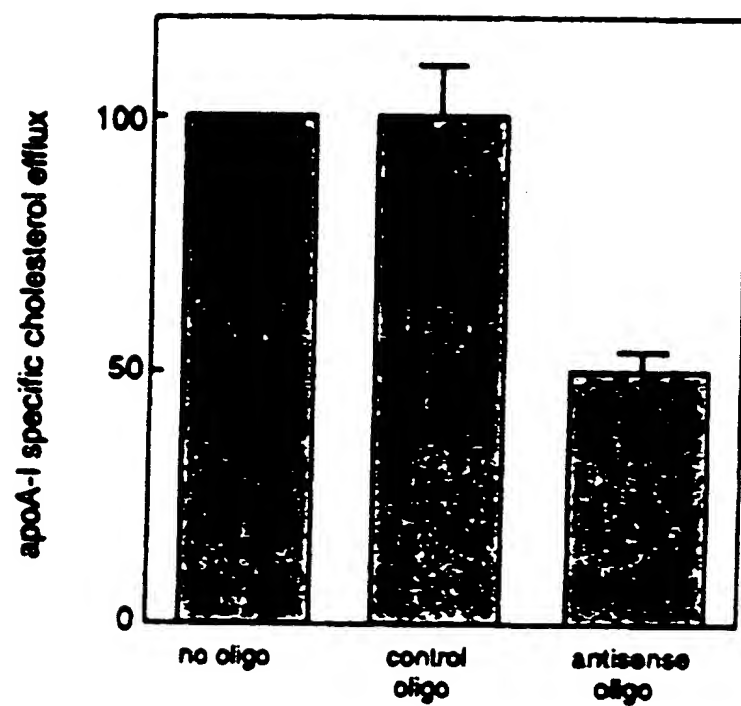


FIG. 6

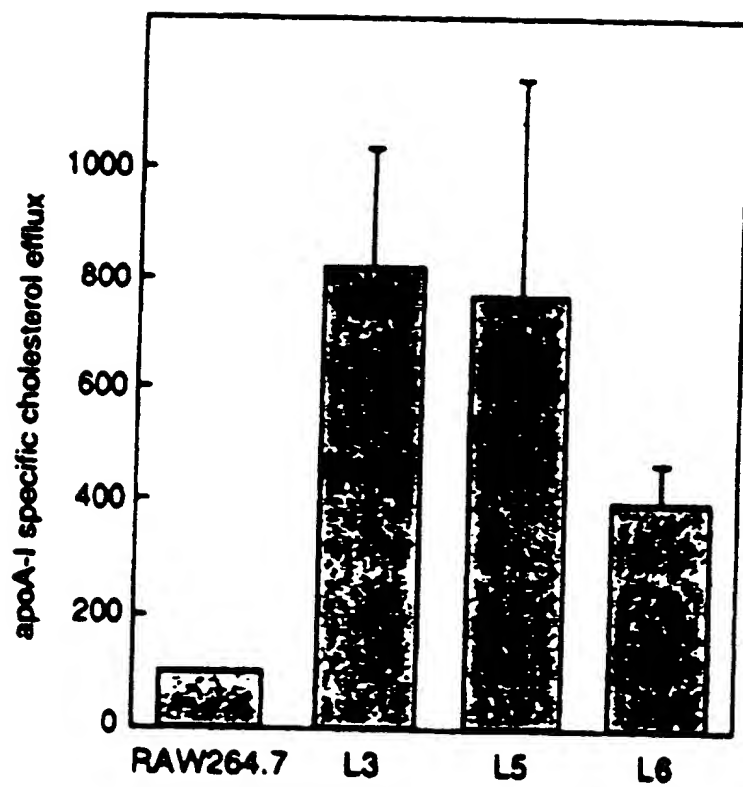


FIG. 7

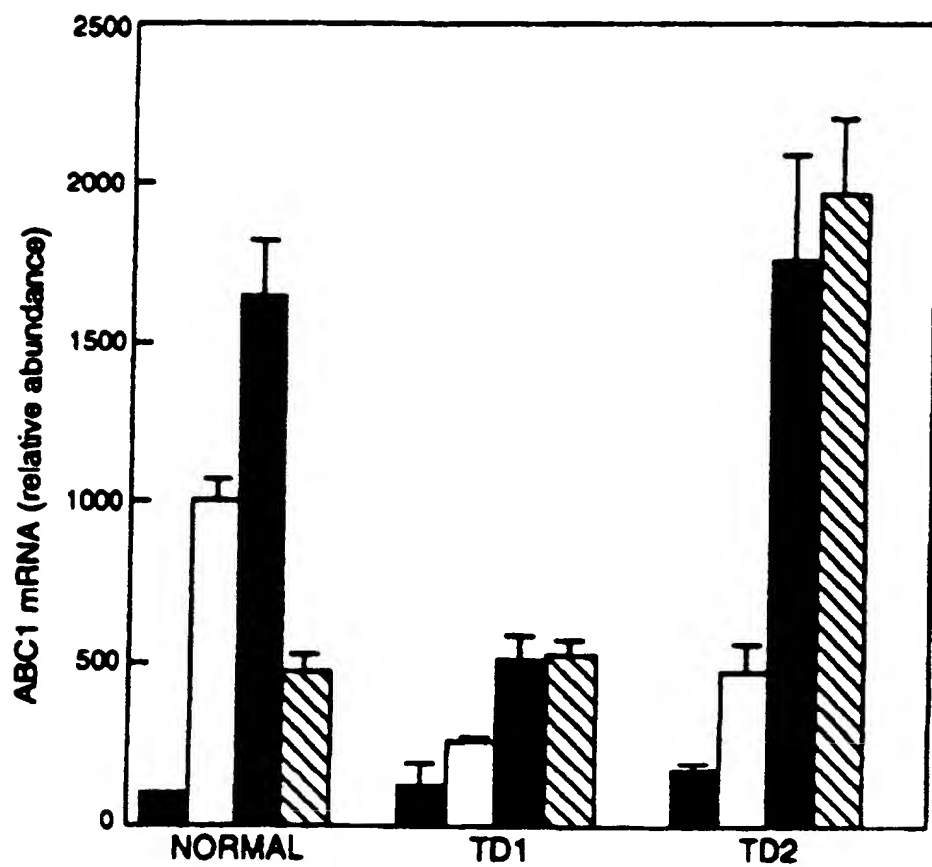


FIG. 8

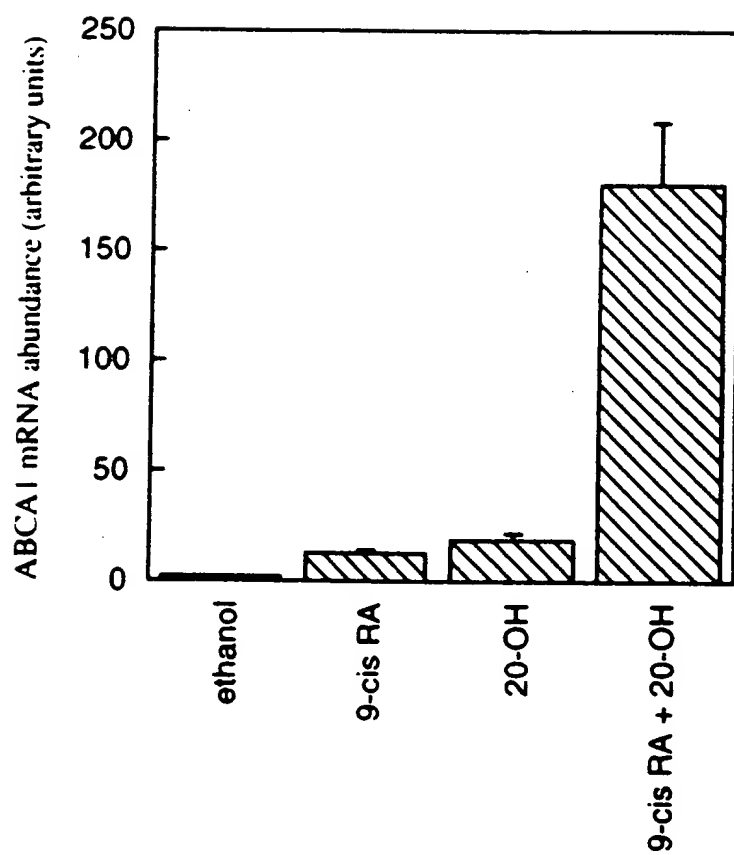


FIG. 9

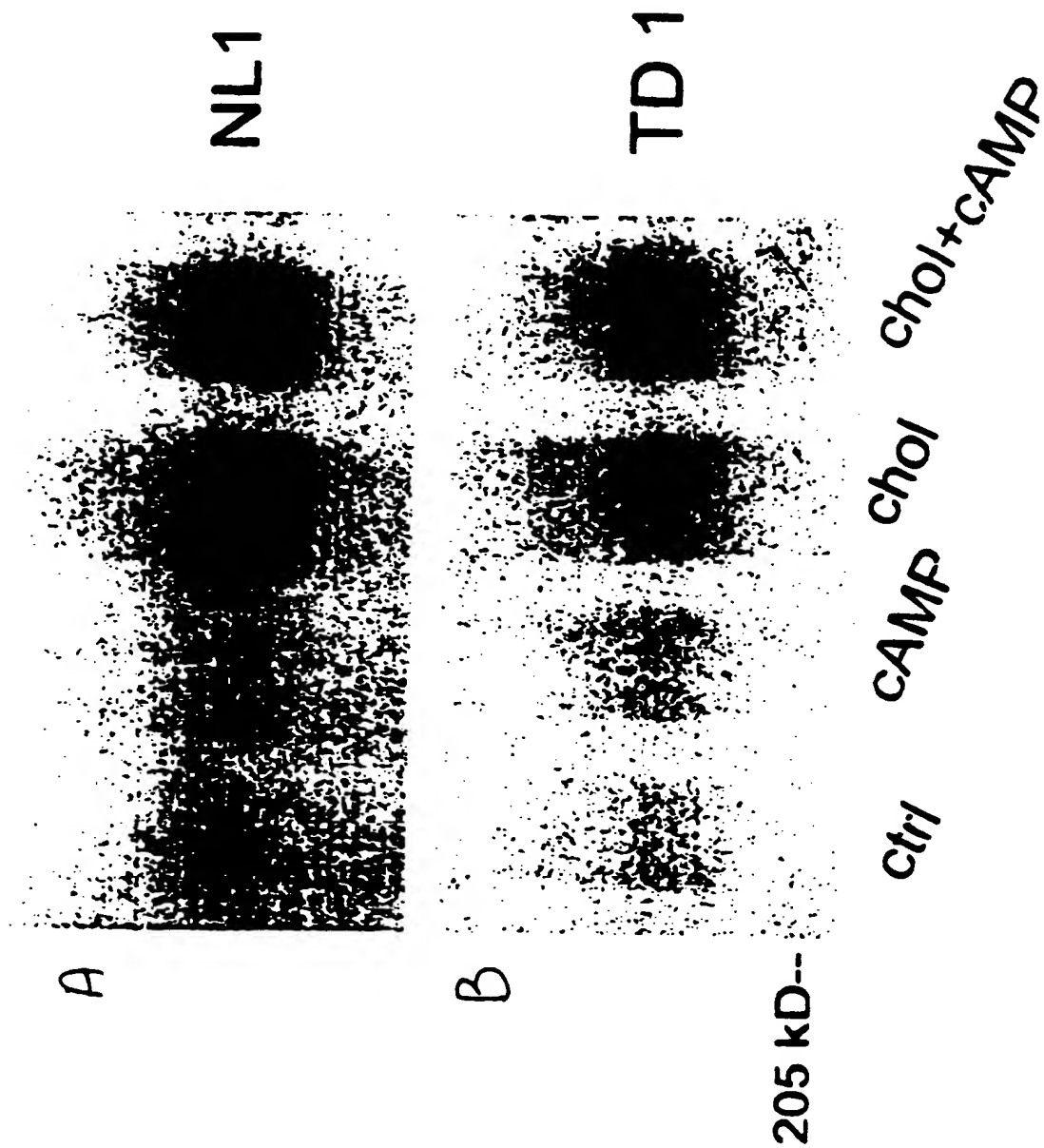


FIG. 10

09596141-061600

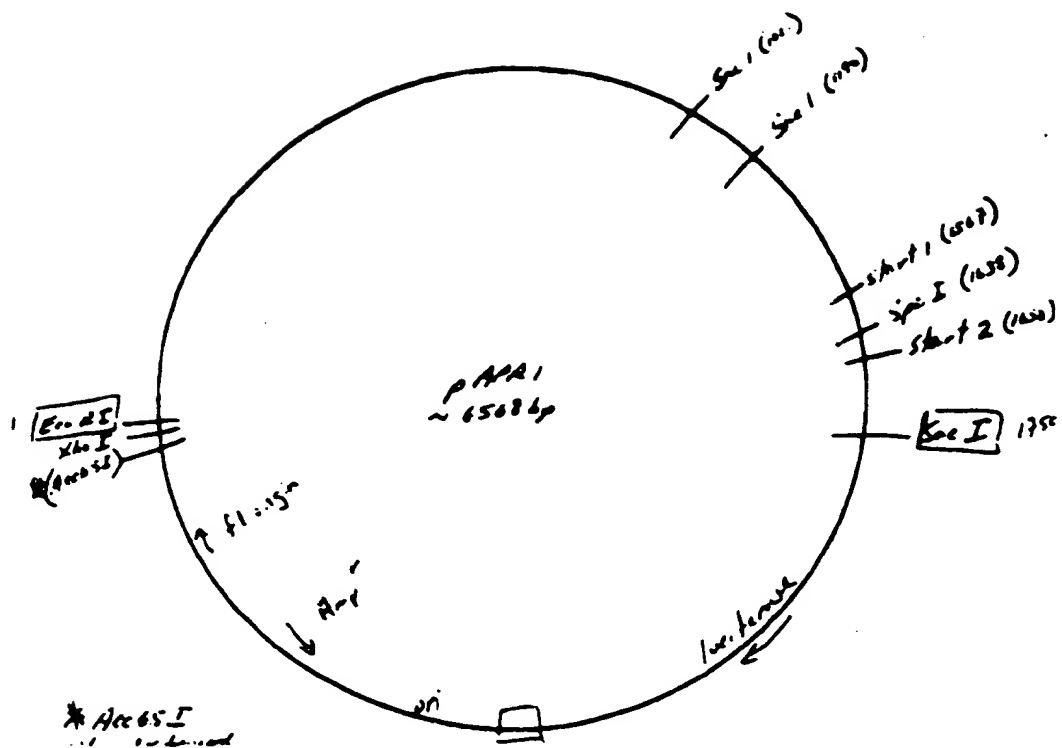


FIG. 11

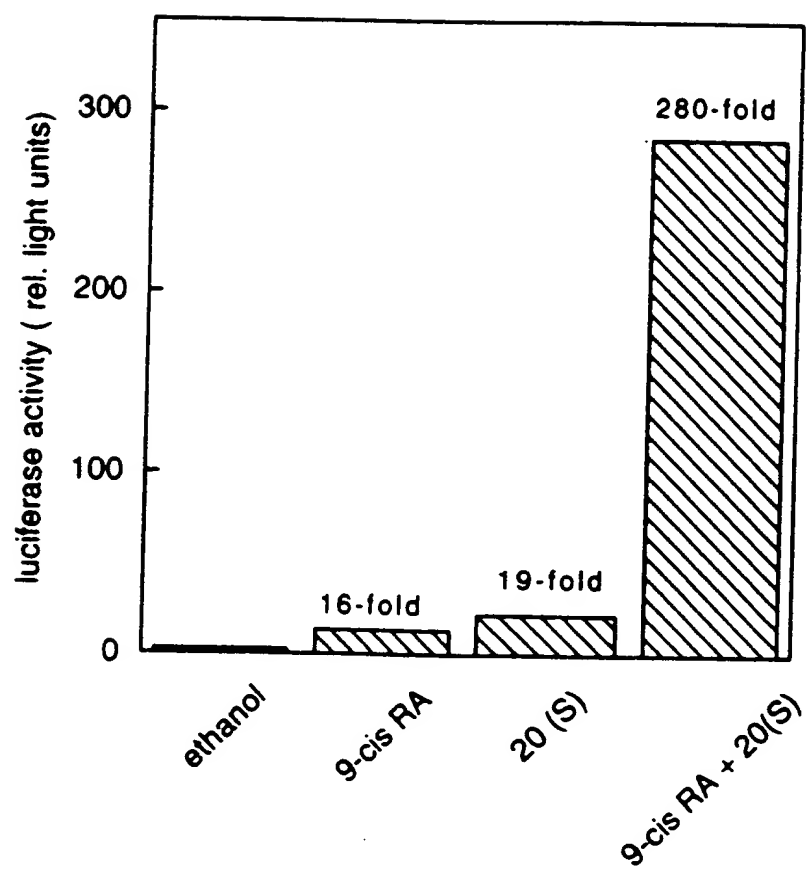


FIG. 12

# ABC1 gene 5' flanking region

1 GAATTCCTTGGTGGCTCCACATGCACATTCAGGGCCCTGCTGGCTCTTCTATAGGGTCTGTCTGAGTGTGATAGAACCACTGATGTGAGTACCTGG  
101 GCTTGAGCGCTGGCCCTGGAGATCCTGTTGACTGTAGCATGGAGGGGCTTGTCAAGCTGAATGTCTGTATGCAGGTGGTGGGAGTTCCTGGAATATGATGGAG  
201 CTGGAGTGGCAAGAGAGTAGGCTTGGGGCAGCTCTCATATGCCACCTCATTTCTGGCCCAAAACTCAGGTCAAACTGTGAAGAGTCTAAATGTGAATCTG  
301 CCCCTTCAAGTGGCTACAAAGGTATCTTTGTCAAGTAGGAGACCTTGTGGCCCTCCACGTGCACCTTCCAGGGCCCTGCTTGGCCCTCTTCTACGGGTCTGTC  
401 CTGAGTCTTCTATGAATCTCCCTTCAGGGCAGATTATATTTAGACTCTTTCACAGTTTGAACCTGAGTTTGGCCCAAGAAATGAAGGTGACATTTAGTTTGTGTTG  
501 GCTTGATGAATGACTTAAATATTTAGACATATGGTGTGTAGGCCCTGCAATTCCTACTCTTGGCCCTTTTTTTTTTGGCCCTCCAGTGTTTTGGGTAGTTTGTGCT  
601 CCCCCCTACAGCCAAAGGCAACAGATAAGTTGGAGGTCTGGAGTGGCTACATAATTTTACAGACTGCAATTTCTCTGGCTGCACCTTCACAAATGTATACATA  
701 AACTAAATACAAAGTCTGTGTTTTTATCACAGGAGGCTGATCAATATAATGAATAATTAAGGGGCTGGTCCCATATTTGTTCTGTGTTTTTGTGTTTT  
801 GTTTCTTTTTTGTGTTTGTGGCCCTCCTTCTCTCAATTTATGAAGAGAAGCAGTAAGATGTTCTCTCGGGTCTCTGAGGGACCTGGGGAGCTCAGGC  
901 TGGGAATCTCCAAGGCAGTAGGTGCGCTATCAAAAATCAAAAGTCCAGGTTTGTGGGGGGAACAAAGCAGCCCCATTACCCAGAGGACTGTCCGCCCTTC  
1001 CCTCACCCAGCCTAGGCCCTTGAAGGAACAAAGACAAAGATGATTTGGCTCTCTGAGGGAGATTTCAGCCCTAGAGCTCTCTCTCCCCCNAATCC  
1101 CTCCCTCCGGCTGAGGAACATAACAAAGGAAAAAATAATTGCGGAAGCAGGATTTAGAGGAAGCAAAATTCACACTGGTGGCCCTTGGCTGCCGGGAACGTG  
1201 GACTAGAGAGTCTGGGGCGCAGCCCCGAGCCCTTCCCGCGTCTTAAGCCCGCGGCCCCCGGGGGAAGGGGACGCAGACCCGGACCCCTAA  
1301 GACACCTGTGTACCTTCCACCCCAACCCCAACCTCCCTAGATGTGTGTGGCGGTGAACGTCGCCCTGTTTAAGGGCGGGGCCCC  
1401 GGCTCCACGTGCTTTCTGCTGAGTGAATGAACATATAAAGAGGCGCGGAACGGGGCGGGGAGGAGGAGAGCACAGGCTTTGACCGATAGTAACCTC  
1501 GCGCTCGGTGCAGCCGAATCTATAAAGGAACCTAGTCCCGGCAAAACCCCGTAATTGCGAGCGAGAGTGAGTGGGGCCGGGACCCGAGAGCCGAGCC  
1601 GACCCCTTCTCCCGGCTGCGGCAGGGCAGGGCGGGGAGCTC

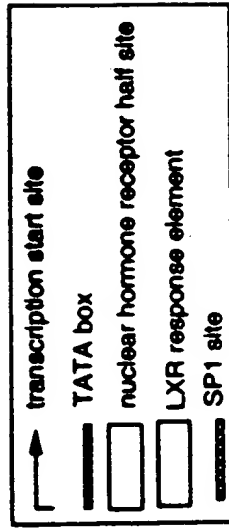


FIG. 13